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Table 1

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Analytical Methods	BLOCKS, HMM, MOTIFS, PRINTS, SPSCAN	BLAST, BLOCKS, HMM, MOTIFS, PFAM, PRINTS, PROFILESCAN	BLAST, BLOCKS, HMM, MOTIFS, PFAM, PRINTS	BLAST, BLOCKS, HMM, MOTIFS, PFAM, PRINTS, PROFILESCAN	BLOCKS, HMM, MOTIFS, PRINTS	BLAST, BLOCKS, HMM, MOTIFS, PRINTS
Identification	Metabotropic glutamate GPCR	Somatostatin-like GPCR	Rhodopsin-like GPCR	Rhodopsin-like GPCR	Metabotropic glutamate GPCR	Secretin-like GPCR
Signature Sequences	M1-A23, I51-V72, G88-P111 C116-A145, I156-L175, M207-P229, G242-T264, E330-K341	I42-V66, P78-M99, W109-I149, V159-L180, T209-L232, V254-T278, Y293-R319	Y44-L74, P62-H83, F109-R131, N143-L164, A231-G255, K278-P304	I46-P70, Y79-I100, L117-F157, R166-S187, S219-F242, L265-L289, S302-K328	I57-L78, G94-E117, C122-V151, L162-L181, M198-F220, G233-L255	N425-T452, I475-W499, A549-L572, F636-N647, Q677-G696, H709-W730
Potential Glycosylation Sites	N191 N405	N13 N16 N23 N58 N84	N8 N110 N300	N7	N30 N352	N88 N110 N127 N281 N392 N424 N443 N505 N647 N785 N798
Potential Phosphorylation Sites	S85 T164 T274 S306 S344 T81 S118 T407 Y312 Y387	S158 T255 S86 T120 S151 S243 S246 T251 T317 S325	T60 T218 S89 S172 T224	S36 S187 T251 S27 T323 S389	S360 S368 S47 T318 S337 S5 T33 S123 T398	T129 S155 S172 S201 S322 S347 S409 S662 S787 S794 S117 T166 T271 T402 T583 T587 T618 S771
Amino Acid Residues	441	353	333	396	403	807
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